(b) a nucleic acid sequence encoding a polypeptide having the amino acid sequence consisting of residues -20-m of SEQ ID NO:2, where m is an integer in the range of +230 to +241;

(c) a nucleic acid sequence encoding a polypeptide having the amino acid sequence consisting of residues n-m of SEQ ID NO:2, where n is an integer in the range of -21 to +64 and m is an integer in the range of +230 to +241;

C1

(d) a nucleic acid sequence encoding a polypeptide consisting of a portion of the complete amino acid sequence encoded by the cDNA clone contained in ATCC Deposit 209023 wherein said portion excludes up to 63 amino acids from the amino terminus and up to 11 amino acids from the C-terminus of said complete amino acid sequence; and

(e) a nucleic acid sequence <u>fully</u> complementary to any of the nucleic acid sequences in (a), (b), (c) or (d), above.

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57. (Once Amended) An isolated polynucleotide comprising a nucleic acid selected from the group consisting of:

C2

(a) a nucleic acid encoding amino acid residues 4 to 63 of SEQ ID NO:2;

(b) a nucleic acid encoding amino acid residues 64 to 242 of SEQ ID NO:2;

(c) a nucleic acid encoding the kringle domain of the polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. 209023;

(d) a nucleic acid encoding the protease domain of the polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. 209023; and

(e) a nucleic acid <u>fully</u> complementary to any of the nucleic acid sequences in (a), (b), (c) or (d), above.

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69. (Once Amended) A method for producing a <u>polypeptide</u> [protein], comprising:

C3